SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT:
 - (A) NAME: Wake Forest University
 - (B) STREET: Medical Center Boulevard
 - (C) CITY: Winston-Salem
 - (D) STATE: North Carolina
 - (E) COUNTRY: USA
 - (F) POSTAL CODE (ZIP): 27157-1023
 - (ii) TITLE OF INVENTION: INACTIVATION OF HIV CO-RECEPTORS AS THERAPY FOR HIV INFECTION
 - (iii) NUMBER OF SEQUENCES: 6
 - (iv) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
 - (vi) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 60/032,277
 - (B) FILING DATE: 02-DEC-1996
- (2) INFORMATION FOR SEQ ID NO / 1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS:
 - (D) TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

Tyr Pro Tyr Asp Val Pro Asp Tyr Ala

- (2) INFORMATION FOR SEQ ID NO: 2:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D)/TOPOLOGY: linear
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

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1

(2)	INFORMATION FOR SEQ ID NO: 3:	•
	(i) SEQUENCE CHARACTERISTICS:	
	(A) LENGTH: 37 base pairs	
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	(D) TOPOLOGY: linear	
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	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
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	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
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	(A) LENGTH: 51 base pairs	
	(B) TYPE: nucleic acid	
	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
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(2)	INFORMATION FOR SEQ ID NO: 6:	
	(i) SEQUENCE CHARACTERISTICS:	
-	(A) LENGTH: 6 amino acids	
	(B) TYPE: amino acid	
	(C) STRANDEDNESS:	
	(D) TOPOLOGY: linear	
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	(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
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<110> CHEN, Si-Yi
     YANG, An-Gang
     BAI, Xuefan
    CHEN, Ji-Dai
     Wake Forest University
<120> INACTIVATION OF HIV CO-RECEPTORS AS THERAPY FOR HIV
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